AND 20 200 THE UNITED STATES PATENT AND TRADEMARK OFFICE

Appl. No.

: 10/734,583

Applicant

: ANDOU et al.

Filed

: December 15, 2003

TC/A.U.

: 3738

Examiner

: Not Yet Assigned

Docket No.

: 2923-595

Customer No.

: 06449

Confirmation No.

: 2600

SUBMISSION OF PRIORITY APPLICATION

Commissioner for Patents P.O. Box 1450 Alexandria, Virginia 22313-1450

Dear Sir:

Submitted herewith is a certified copy of European Patent Application No. 99115613.4 filed August 6, 1999 and Japanese Patent Nos. JP 8/355812 filed December 25, 1996 and JP 10/141379 May 22, 1998, from which priority has been claimed in the above-referenced patent application.

Respectfully submitted,

Rv

Steven M. Giovannetti Attorney for Applicants Registration No. 51,739

ROTHWELL, FIGG, ERNST & MANBECK, p.c.

Suite 800, 1425 K Street, N.W.

Washington, D.C. 20005 Telephone: (202)783-6040

3 m m

THIS PAGE BLANK (USPTO)



Europäisches **Patentamt**

European **Patent Office**

Office européen des brevets

Bescheinigung

Certificate

Attestation

Die angehefteten Unterlagen stimmen mit der ursprünglich eingereichten Fassung der auf dem nächsten Blatt bezeichneten europäischen Patentanmeldung überein.

The attached documents are exact copies of the European patent application conformes à la version described on the following page, as originally filed.

Les documents fixés à cette attestation sont initialement déposée de la demande de brevet européen spécifiée à la page suivante.

Patentanmeldung Nr.

Patent application No. Demande de brevet n°

99115613.4

CERTIFIED COPY OF PRIORITY DOCUMENT

Der Präsident des Europäischen Patentamts; Im Auftrag

For the President of the European Patent Office

Le Président de l'Office européen des brevets p.o.

I.L.C. HATTEN-HECKMAN

DEN HAAG, DEN THE HAGUE, LA HAYE, LE

24/08/00

THIS PAGE BLANK (USPTO)



Europäisches Patentamt

European **Patent Office** Office européen des brevets

Blatt 2 der Bescheinigung Sheet 2 of the certificate Page 2 de l'attestation

Anmeldung Nr.:

Application no.: Demande n°:

99115613.4

Anmeldetag: Date of filing: Date de dépôt:

06/08/99

An melder: Applicant(s): Demandeur(s):

HyGene AG, c/o Mäder + Baumgartner Treuhand AG

8121 Neuhausen am Rheinfall

SWITZERLAND

Bezeichnung der Erfindung: Title of the invention: Titre de l'invention:

Monomeric protein of the TGF-beta family

÷:

In Anspruch genommene Prioriät(en) / Priority(ies) claimed / Priorité(s) revendiquée(s)

Staat:

State:

Tag:

Aktenzeichen:

File no.

Pays:

Date: Date:

Numéro de dépôt:

Internationale Patentklassifikation: International Patent classification: Classification internationale des brevets:

C12N15/12, C07K14/495, A61K38/18

Am Anmeldetag benannte Vertragstaaten:
Contracting states designated at date of filing: AT/BE/CH/CY/DE/DK/ES/FI/FR/GB/GR/IE/IT/LI/LU/MC/NL/PT/SE Etats contractants désignés lors du depôt:

Bemerkungen: Remarks: Remarques:

"HIS PAGE BLANK (USPTO)

Ex. 1

DESC

PATENTANWÄLTE

European Patent Attorneys
European Trade Mark Attorneys

DIPL.-ING. H. WEICKMANN
DIPL.-ING. F. A. WEICKMANN
DIPL.-CHEM. B. HUBER
DR.-ING. H. LISKA
DIPL.-PHYS. DR. J. PRECHTEL
DIPL.-CHEM. DR. B. BÖHM
DIPL.-CHEM. DR. W. WEISS
DIPL.-PHYS. DR. J. TIESMEYER
DIPL.-PHYS. DR. M. HERZOG
DIPL.-PHYS. B. RUTTENSPERGER

POSTFACH 860 820 81635 MÜNCHEN

KOPERNIKUSSTRASSE 9 81679 MÜNCHEN

TELEFON (089) 45563 0
TELEX 522 621
TELEFAX (089) 45563 999
E-MAIL email@weickmann.de

Unser Zeichen: 20780P EP/BBcl

Anmelder:
HyGene AG
c/o Mäder + Baumgartner
Treuhand AG
Schwanenfelsstrasse 10a
8121 Neuhausen am Rheinfall

Schweiz

EPO-Munich 52 **06. Aug. 1999**

Monomeric Protein of the TGF-B Family

- 1 -

EPO-Munich 52

06. Aug. 1999

Monomeric Protein of the TGF-ß Family

Description

5

10

15

The present invention concerns a biologically active protein from the TGF-ß superfamily, wherein this protein remains in monomeric form due to substitution or deletion of a cysteine which is reponsible for the dimerization in the wild-type protein. Further the invention concerns a nucleic acid, which codes for a protein according to the invention, an expression vector containing such nucleic acid and a host cell, containing a corresponding nucleic acid or an expression vector, said nucleic acid being suitable for the expression of the protein. The invention also concerns a pharmaceutical composition containing the protein according to the invention or a nucleic acid coding therefor. The use of the pharmaceutical composition according to the invention concerns the prevention or treatment of all conditions which can also be treated with the dimeric form of the corresponding protein.

20 Man
Deveroreles
whice
form
25 factor
95 (

Many growth factors from the TGF-β superfamily (Kingsley, Genes and Development 8, 133-146 (1994) as well as the references cited therein) are relevant for a wide range of medical treatment methods and applications which in particular concern promotion of cell proliferation and tissue formation, including wound healing and tissue reproduction. Such growth factors in particular comprise members of the TGF-β (transforming growth factor, cf. e.g. Roberts and Sporn, Handbook of Experimental Pharmacology 95 (1990), page 419-472, editors: Sporn and Roberts), the DVR-group (Hötten et al., Biochem. Biophys. Res. Comm. 206 (1995), page 608-613 and further literature cited therein) including BMPs (bone morphogenetic protein, cf. e.g. Rosen and Thies, Growth Factors in Perinatal Development (1993), page 39-58, editors: Tsang, Lemons and Balistreri) and GDFs (growth differentiation factors), the inhibin/activin (cf. e.g. Vale et al., The

30

10

15

20

25

30

- 2 -

Physiology of Reproduction, second edition (1994), page 1861-1878, editors: Knobil and Neill) and the GDNF protein family (Rosenthal, Neuron 22 (1999), page 201-203; Airaksinen et al. Mol Cell Neurosci 13 (1999), page 313-325). Although the members of the TGF-ß superfamily show high amino acid homologies in the mature part of the protein, in particular 7 conserved cysteines, they show considerable variations in their exact functions. Often individual growth factors of these families exhibit a plurality of functions at the same time, so that their application is of interest in various medical indications. Some of these multifunctional proteins also have survival promoting effects on neurons in addition to functions such as e.g. regulation of the proliferation and differention in many cell types (Roberts and Sporn, supra; Sakurai et al., J. Biol. Chem. 269 (1994), page 14118-14122). Thus e.g. trophic effects on embryonic motoric and sensory neurons were demonstrated for TGF-\$\textit{\beta}\$ in vitro (Martinou et al., Devl. Brain Res. 52, page 175-181 (1990) and Chalazonitis et al., Dev. Biol. 152, page 121-132 (1992)). In addition, effects promoting survival are shown for dopaminergic neurons of the mid-brain for the proteins TGF-\$\beta-1\$, -2, -3, activin A and GDNF (glial cell line-derived neurotrophic factor), a protein which has structural similarities to TGF-B superfamily members, these effects being not mediated via astrocytes (Krieglstein et al., EMBO J. 14, page 736-742 (1995)).

Interesting members of the TGF-ß superfamily or active variants thereof comprise the TGF-ß proteins like TGF-ß1, TGF-ß2, TGF-ß3, TGF-ß4, TGF-ß5 (U.S. 5,284,763; EP 0376785; U.S. 4,886,747; DNA 7 (1988), page 1-8), EMBO J. 7 (1988), page 3737-3743), Mol. Endo. 2 (1988), page 1186-1195), J. Biol. Chem. 265 (1990), page 1089-1093), OP1, OP2 and OP3 proteins (U.S. 5,011,691, U.S. 5,652,337, WO 91/05802) as well as BMP2, BMP3, BMP4 (WO 88/00205, U.S. 5,013,649 and WO 89/10409, Science 242 (1988), page 1528-1534), BMP5, BMP6 and BMP-7 (OP1) (Proc. Natl. Acad. Sci. 87 (1990), page 9841-9847, WO 90/11366), BMP8 (OP2) (WO 91/18098), BMP9 (WO 93/00432), BMP10 (WO 94/26893),

10

15

20

25

30

- 3 -

BMP11 (WO 94/26892), BMP12 (WO 95/16035), BMP13 (WO95/16035), BMP15 (WO 96/36710), BMP16 (WO 98/12322), BMP3b (Biochem. Biophys. Res. Comm. 219 (1996), page 656-662), GDF1 (WO 92/00382 and Proc. Natl. Acad. Sci. 88 (1991), page 4250-4254), GDF8 (WO 94/21681), GDF10 (WO95/10539), GDF11 (WO 96/01845), GDF5 (CDMP1, MP52) (WO 95/04819; WO96/01316; WO 94/15949, WO 96/14335 and WO 93/16099 and Nature 368 (1994), page 639-643), GDF6 (CDMP2, BMP13) (WO 95/01801, WO 96/14335 and WO95/16035), GDF7 (CDMP3, BMP12) (WO 95/01802 and WO 95/10635), GDF14 (WO 97/36926), GFD15 (WO 99/06445), GDF16 (WO 99/06556), 60A (Proc.Natl. Acad. Sci. 88 (1991), page 9214-9218), DPP (Nature 325 (1987), page 81-84), Vgr-1 (Proc. Natl. Acad. Sci. 86 (1989), page 4554-4558) Vg-1, (Cell 51 (1987), page 861-867), dorsalin (Cell 73 (1993), page 687-702), MIS (Cell 45 (1986), page 685-698), pCL13 (WO 97/00958), BIP (WO 94/01557), inhibin a, activin ßA and activin ßB (EP 0222491), activin &C (MP121) (WO 96/01316), activin &E and GDF12 (WO 96/02559 and WO 98/22492), activin ßD (Biochem. Biophys. Res. Comm. 210 (1995), page 581-588), GDNF (Science 260 (1993), page 1130-1132, WO 93/06116), Neurturin (Nature 384 (1996), page 467-470), Persephin (Neuron 20 (1998), page 245-253, WO 97/33911), Artemin (Neuron 21 (1998), page 1291-1302), Mic-1 (Proc. Natl. Acad. Sci USA 94 (1997), page 11514-11519), Univin (Dev. Biol. 166 (1994), page 149-158), ADMP (Development 121 (1995), page 4293-4301), Nodal (Nature 361 (1993), page 543-547), Screw (Genes Dev. 8 (1994), page 2588-2601). Other useful proteins include biologically active biosynthetic constructs including biosynthetic proteins designed using sequences from two or more known morphogenetic proteins. Examples of biosynthetic constructs are disclosed in U.S. 5,011,691 (e.g. COP-1, COP-3, COP-4, COP-5, COP-7 and COP-16). The disclosure of the cited publications including patents or patent applications are incorporated herein by reference.

10

15

- 4 -

The occurence of proteins of the TGF- β superfamily in various tissuous stages and development stages corresponds with differences with regard to their exact functions as well as target sites, life span, requirements for auxiliary factors, necessary cellular physiological environment and/or resistance to degradation.

The proteins of the TGF- β superfamily exist as homodimers or heterodimers having a single disulfide bond. This disulfide bond is mediated by a specific and in most of the proteins conserved cysteine residue of the respective monomers. Up to now it was considered as indispensible for the biological activity that the protein is present in its dimeric form. Several publications indicated that biological activity can only be obtained for dimeric proteins and it was speculated that this dimer formation is important for further polymer formation of two or more dimers to achieve intercellular signal transmission by simultaneous binding to type I and type II receptors for the TGF- β superfamily proteins on cells. It was assumed that only this simultaneous binding to both kinds of receptors would allow for effective intercellular signal transmission for the benefit of the patient (Bone, volume 19 (1996), page 569-574).

20

A disadvantage of the use of these proteins as medicaments and their production is, that they are not readily obtainable in biologically active and sufficiently pure form by recombinant expression in prokaryots without intensive renaturation procedures.

25

Thus it was the object of the present invention to provide a simple and inexpensive possibility to reproducibly produce proteins exhibiting high biological activity, wherein this biological activity should essentially correspond to that of the dimers of the proteins of said families.

30

This object is solved according to the invention by a protein selected from the members of the TGF-ß protein superfamily, such protein being

10

15

20

- 5 -

necessarily monomeric due to substitution or deletion of a cysteine which is responsible for dimeric formation.

Surprisingly it has been found that the substitution or deletion of the cysteine, which normally effects the dimerization in the proteins, results upon expression and correct folding (proper formation of the intramolecular disulfide bridges) in a monomeric protein that retains the biological activity of the dimeric form. Even more surprisingly, it was found that at least some of the monomeric proteins show a higher activity, based on the weight of protein, than their respective dimeric forms. Apart from this improved biological activity an important advantage for the proteins according to the invention is that they can be expressed in a large amount in prokaryotic hosts and upon simple refolding of the monomers they are obtained in high purity and very high yield without the need to separate dimerized from nondimerized (monomeric) protein. The findings of the present invention are very surprising since, as already mentioned above, it was common understanding that only a dimer of the morphogenetic proteins has biological activity. Despite this understanding the proteins according to the invention show an up to two-fold higher activity than that of the dimer on the basis of protein weight. The smaller size of the proteins of the invention, while maintaining the biological activity, can also be considered as advantageous, e.g. for applications concerning the brain since the monomeric protein can much easier pass the blood-brain-barrier than the dimeric form.

25

30

The proteins according to the invention encompass all proteins of the mentioned protein families that are normally present in dimeric form. Also parts of such proteins that retain substantial activity or fusion proteins or precursor forms of proteins shall be considered as encompassed by the present invention as well as biologically active naturally occurring or biosynthetic variants of TGF-ß superfamily proteins, as long as they show at least considerable biological activity.

10

15

- 6 -

In a preferred embodiment of the present invention the monomeric protein is a mature protein or a biologically active part or variant thereof. The term "biologically active part or variant thereof" is meant to define either fragments retaining activity, precursor proteins that are e.g. cleaved at the site of activity to the mature form or show biological activity themselves, or also variants that still maintain essentially the biological activity of the wild-type protein. Such variants preferably contain conservative amino acid substitutions, but especially at the N-terminal part of the mature proteins even considerable deletions or substitutions do not lead to a considerable loss of biological activity. It is well within the skill of the man in the art to determine whether a certain protein shows the required biological activity. Proteins showing at least 70% and preferably at least 80% homology to the mature wild-type proteins of the above referenced protein families should be understood as encompassed by the present invention, as long as they contain the deletion or substitution of a cysteine, as required for the proteins according the invention, and therefore do not form dimers.

It is especially preferred that proteins according to the invention contain at least the 7 cysteine region characteristic for the TGF- β protein superfamily.

20

This specific 7 cysteine region is considered to be the most important part of the proteins in view of the biological activity. Therefore proteins retaining this critical region are preferred proteins according to the invention. It is disclosed in the state of the art which cysteine is responsible in a certain protein family or protein for dimer formation (see for example: Schlunegger & Grutter (1992) Nature 358, 430-434; Daopin et al., (1992) Science 257, 369-373 and Griffith et at., Proc. Natl. Acad. Sci. 93 (1996), page 878-883). This cysteine has to be deleted or substituted by another amino acid to form a protein according to the invention.

30

25

The 7 cysteine region is known for many proteins of the TGF- β protein superfamily. In this region the respective location of the cysteine residues

15

20

25

30

-7-

to each other is important and is only allowed to vary slightly in order not to lose the biological activity. Consensus sequences for such proteins are known in the state of the art and all proteins complying with such consensus sequences are considered to be encompassed by the present invention.

In an especially preferred embodiment of the present invention the protein contains a consensus sequence according to the following sequence

wherein C denotes cysteine, Y denotes any amino acid including cysteine and X denotes any amino acid except cysteine.

More preferably the protein according to the invention contains a consensus sequence according to the following sequence

C (Y)₂₈ C Y Y Y C (Y)₃₀₋₃₂ X C (Y)₃₁ C Y C (Formula II), wherein C, X and Y have the same meaning as defined above.

Even more preferably the protein according to the invention contains a consensus sequence according to the following sequence

C (X) $_{28}$ C X X X C (X) $_{31-33}$ C (X) $_{31}$ C X C (Formula III), wherein C and X have the same meaning as defined above.

In these consensus sequences especially preferred distances between the respective cysteine residues are contained, wherein also already the dimer forming cysteine is substituted by another amino acid. As with all proteins of said protein superfamily the location of and distance between the cysteines is more important than the identity of the other amino acids contained in this region. Therefore, the consensus sequence shows the respective location of the cysteines, but does not show the identity of the

10

15

20

25

30

-8-

other amino acids, since these other amino acids are widely variable in the proteins of the TGF- β protein superfamily.

In a preferred embodiment of the present invention the monomeric protein according to the invention is a morphogenetic protein.

Most of the members of the TGF-ß protein superfamily are morphogenetic proteins that are useful for treatments where regulation of differentiation and proliferation of cells or progenitor cells is of interest. This can result in replacement of damaged and/or diseased tissue like for example skeletal (bone, cartilage) tissue, connective tissue, periodontal or dental tissue, neural tissue, tissue of the sensory system, liver, pancreas, cardiac, blood vessel and renal tissue, uterine or thyroid tissue etc. Morphogenetic proteins are often useful for the treatment of ulcerative or inflammatory tissue damage and wound healing of any kind such as enhanced healing of ulcers, burns, injuries or skin grafts. Especially preferred proteins according to the invention belong to the TGF- β , BMP, GDF, activin or GDNF families. Several BMP proteins which were originally discovered by their ability to induce bone formation, have been described, as also indicated above. Meanwhile, several additional functions have been found as it is also true for members of the GDFs. These proteins show a very broad field of applications and especially are in addition to their bone and cartilage growth promoting activity (see for example: WO 88/00205, WO 90/11366, WO 91/05802) useful in periodontal disease, for inhibiting periodontal and tooth tissue loss, for sealing tooth cavities, for enhancing integration of a tooth in a tooth socket (see for example: WO 96/26737, WO 94/06399, WO 95/24210), for connective tissue such as tendon or ligament (see for example: WO 95/16035), for improving survival of neural cells, for inducing growth of neural cells and repairing neural defects, for damaged CNS tissue due to stroke or trauma (see for example: WO 97/34626, WO 94/03200, WO 95/05846), for maintaining or restoring sensory perception (see for example WO 98/20890, WO 98/20889), for renal failure (see for example:

15

20

25

30

- 9 -

WO 97/41880, WO 97/41881), for liver regeneration (see for example WO 94/06449), for regeneration of myocardium (see for example WO 98/27995), for treatment or preservation of tissues or cells for organ or tissue transplantation, for integrity of gastrointestinal lining (see for example WO 94/06420), for increasing progenitor cell population as for example hematopoietic progenitor cells by ex vivo stimulation (see for example WO 92/15323), etc. One preferred member of the GDF family is the protein MP52 which is also termed GDF-5 or CDMP-1. Applications for MP52 reflect several of the already described applications for the BMP/GDF family. MP52 is considered to be a very effective promoter of bone and cartilage formation as well as connective tissue formation (see for example WO 95/04819, Hötten et al., (1996), Growth Factors 13, 65-74, Storm et al., (1994) Nature 368, 639-643, Chang et al., (1994) J. Biol. Chem. 269 (45), 28227-28234). In this connection MP52 is useful for applications concerning the joints between skeletal elements (see for example Storm & Kingsley (1996) Development 122, 3969-3979). One example for connective tissue is tendon and ligament (Wolfman et al., (1997), J. Clin. Invest. 100, 321-330, Aspenberg & Forslund (1999), Acta Orthop Scand 70, 51-54, WO 95/16035). MP52 is also useful for tooth (dental and periodontal) applications (see for example WO 95/04819, WO 93/16099, Morotome et al. (1998), Biochem Biophys Res Comm 244, 85-90). MP52 is useful in wound repair of any kind. It is in addition very useful for promoting tissue growth in the neuronal system and survival of dopaminergic neurons, for example. MP52 in this connection is useful for applications in neurodegenerative diseases like e.g. Parkinson's disease and possibly also Alzheimer's disease for Huntington chorea tissues (see for example WO 97/03188, Krieglstein et al., (1995) J. Neurosci Res. 42, 724-732, Sullivan et al., (1997) Neurosci Lett 233, 73-76, Sullivan et al. (1998), Eur. J. Neurosci 10, 3681-3688). MP52 allows to maintain nervous function or to retain nervous function in already damaged tissues. MP52 is therefore considered to be a generally applicable neurotrophic factor. It is also useful for diseases of the eye, in particular retina cornea and optic

10

15

20

25

30

- 10 -

nerve (see for example WO 97/03188, You et al. (1999), Invest Opthalmol Vis Sci 40, 296-311). The monomeric MP52 is expected to show all the already described activities of the dimeric form as well as some further described activities as described for the dimeric BMP/GDF family members. It is expected to be for example also useful for increasing progenitor cell populations and for stimulating differentiation of progenitor cells ex vivo. Progenitor cells can be cells which take part in the cartilage formation process or hematopoietic progenitor cells. It is also useful for damaged or diseased tissue where a stimulation of angiogenesis is advantageous (see for example: Yamashita et al. (1997), Exp Cell Res 235, 218-226).

An especially preferred protein according to the invention therefore is protein MP52 or a biologically active part or variant thereof. Like in the already above mentioned definition of these terms MP52 can e.g. be used in its mature form, however, it can also be used as a fragment thereof at least containing the 7 cysteine region or also in a precursory form. Deviations at the N-terminal part of mature MP52 do not affect its activity to a considerable degree. Therefore, substitutions, deletions or additions on the N-terminal part of the proteins are still within the scope of the present invention. It might be useful to add a peptide to the N-terminal part of the protein, e.g. for purification reasons. It might not be necessary to cleave off this added peptide after expression and purification of the protein. Additional peptides at the N- or C-terminal part of the protein may also serve for the targeting of the protein to special tissues such as nerve or bone tissue or for the penetration of the blood/brain barrier. Generally, also fusion proteins of a monomeric protein according to the invention and another peptide or group are considered within the scope of the present invention, wherein these other peptides or groups are directing the localization of the fusion protein, e.g. because of an affinity to a certain tissue type etc. Examples for such fusion proteins are described in WO 97/23612. The protein containing such addition will retain its biological

10

15

20

25

30

- 11 -

activity at least as long as such addition does not impair the formation of the biologically active conformation of the protein.

In an especially preferred embodiment of the present invention the proteins comprises the amino acid sequence according to SEQ.ID.NO.1 (DNA and protein sequence) and SEQ.ID.No.2 (protein sequence, only), respectively. SEQ.ID.NO.2 shows the complete protein sequence of the prepro protein of human MP52, as already disclosed in WO 95/04819. The start of the mature protein lies preferably in the area of amino acids 352-400, especially preferred at amino acids 381 or 382. Therefore, the mature protein comprises amino acids 381-501 or 382-501. The first alanine of the mature protein can be deleted and the mature protein then preferably comprises amino acids 383-501. The cysteine at position 465 that is present in the already described dimeric MP52 protein is according to the invention either deleted or substituted by another amino acid. This deletion or substitution is represented by Xaa at the respective position in SEQ.ID.Nos.1 and 2.

The activin/inhibin family proteins are of interest for applications related to contraception, fertility and pregnancy (see for example WO 94/19455, U.S. 5,102,868). They are also of interest for applications like repair or prevention of diseases of the nervous system, they can be used in the repair of organ tissue such as liver and even in bone and cartilage, too. In this connection MP121 (activin &C) is especially useful in applications for growth or regeneration of damaged and/or diseased tissue, especially the liver tissue, neural tissue, skeletal tissue (see for example WO 96/01316, WO 98/22492 and WO 97/03188). MP121 is known to be predominantly expressed in the liver whereby the mRNA is markedly reduced after partial hepatectomy. MP121 is expected to regulate the liver mass (Zhang et al., Endocrine Journal 44 (1997), page 759-764). The monomeric MP121 shows all the already described activities of the dimeric form as well as some further described activities as described for the dimeric TGF-& superfamily members. It is for example also expected to be useful in

- 12 -

treatment of ulceration (for example stomach ulceration) and useful for integrity of gastrointestinal lining and for stimulating differentiation of progenitor cells <u>ex vivo</u>, treatment or preservation of mammalian tissue or cells, e.g. for organ or tissue transplantation.

5

10

15

A further preferred protein according to the invention therefore is MP121, a member of the activin/inhibin protein family. Also for this protein a biologically active part or variant thereof is encompassed by the present invention according to the above defined rules. An especially preferred embodiment is shown in SEQ.ID.NO.3 (DNA and protein sequence) and SEQ.ID.NO.4 (protein sequence, only) respectively. SEQ.ID.NO.4 shows the complete amino acid sequence of the prepro protein of human MP121, that has already been disclosed in WO 96/01316. The start of the mature protein lies preferably between amino acids 217 and 247, most preferred at amino acid 237. A preferred mature protein therefore comprises the mature part of the protein starting at amino acid 237 and ending at amino acid 352. However, also the precursor protein comprising the whole shown amino acid sequence is encompassed by the present invention. The cysteine at position 316 is according to the invention either deleted or substituted by another amino acid, being represented by Xaa in SEQ.ID.Nos.3 and 4.

20

The amino acid by which the cysteine residue effecting the dimerization is substitued can be selected by any amino acid that does not impair the formation of a biologically active conformation. The amino acid is preferably selected from the group of alanine, serine, threonine, leucine, isoleucine, glycine and valine.

25

30

The proteins according to the invention are in summary characterized by the absence of the cysteine residue in the amino acid sequence responsible for the dimer formation. This absence can be effected by substitution of this cysteine by another amino acid or by deletion. In case of deletion, however, it must be assured for the protein that the formation of the biologically

20

25

30

- 13 -

active conformation is not hindered. The same is true for the selection of the substitution amino acid, wherein it is preferred to use an amino acid which has a form similar to cysteine.

The monomeric proteins according to the invention can be easily produced, in particular by expression in prokaryots and renaturation according to known methods. It is advantageous that the protein can be obtained in exceedingly biologically active form. The proteins exhibit in monomeric form about the same activity as the dimer so that based on the amount of active substance only half of the monomeric protein has to be used in order to obtain the same positive biological effects.

A further subject matter of the present invention is a nucleic acid encoding a protein according to the invention. It is obvious that the nucleic acid has to have such a sequence that a deletion or substitution of the cysteine responsible for the dimer formation is achieved. The nucleic acid can be a naturally occurring nucleic acid, but also a recombinantly produced or processed nucleic acid. The nucleic acid can be both a DNA sequence and an RNA sequence, as long as the protein according to the invention can be obtained from this nucleic acid upon expression in a suitable system.

In a preferred embodiment of the invention the nucleic acid is a DNA sequence. This DNA sequence in an especially preferred embodiment of the invention comprises a sequence as shown in SEQ.ID.NO.1 and SEQ.ID.NO.3, respectively, or parts thereof. SEQ.ID.NO.1 shows a nucleic acid encoding MP52, wherein the codon for the cysteine responsible for the dimer formation is replaced by another codon which does not encode cysteine or deleted. This substitution or deletion is shown as "nnn" in the sequence protocols. SEQ.ID.NO.3 shows a nucleic acid encoding MP121, wherein also the codon for the cysteine effecting the dimer formation is replaced by a respective different codon or deleted. Instead of the complete

10

15

- 14 -

sequences of SEQ.ID.NOs.1 or 3 also parts can be used that encode the mature proteins or fragments also described above.

It is preferred in the framework of the present invention that the nucleic acid apart from the coding sequences also contains expression control sequences. Such expression control sequences are known to the man skilled in the art and serve to control the expression of the encoded protein in a host cell. The host cell does not have to be an isolated cell, moreover, the nucleic acid can be expressed in the patient in vivo in the target tissue. This can be done by inserting the nucleic acid into the cell genome, however, it is also possible to transform host cells with expression vectors containing a nucleic acid according to the invention. Such expression vectors are a further subject matter of the present invention, wherein the nucleic acid is inserted in a suitable vector system, the vector system being selected according to the desired expression of the protein. The vector system can be a eukaryotic vector system, but - in the framework of the present invention - it is preferably a prokaryotic vector system, with which the proteins can be produced in prokaryotic host cells in a particularly easy and pure manner. In addition, the expression vector can be a viral vector.

20

25

Also host cells in turn are a further subject matter of the present invention. The host cells are characterized in that they contain a nucleic acid according to the invention or an expression vector according to the invention and that they are able to use the information present in the nucleic acids and in the expression vector, respectively, for the expression of a monomeric protein according to the invention.

30

Although in the framework of the present invention also eukaryotic host cells are suitable for the production of the protein, it is, as mentioned already several times above, particularly advantageous that the protein according to the invention can be produced in prokaryotic host cells, which therefore represent a preferred embodiment of the present invention.

15

20

25

30

- 15 -

After such preferred expression in prokaryotic host cells the protein is purified and renatured according to known methods, thereby effecting intramolecular cystine bridge formation.

Since, however, not only in vitro production of the monomeric protein is possible, but also in vivo expression of a nucleic acid according to the invention, a further preferred embodiment is a eukaryotic host cell, and especially a eukaryotic host cell containing the DNA in its genome, or as an expression vector. Such host cell can also be useful for application to an individual in need of morphogenic treatment. 10

Further subject matters of the present application are pharmaceutical compositions comprising at least one monomeric protein according to the invention or at least one nucleic acid encoding for such a protein or at least one corresponding expression vector, or at least one eukaryotic host cell expressing the monomeric protein.

The protein itself, but also a nucleic acid according to the invention, an expression vector or a host cell can be considered to be advantageous as active substances in a pharmaceutical composition. Also combinations of monomeric proteins, with either biological activities in the same or different applications, can be used in preferred pharmaceutical compositions. Especially preferred for neuronal applications are combinations of MP52 with other TGF-\$\beta\$ superfamily proteins, both in monomeric form, like for example with GDNF (see WO 97/03188). Also preferred for neuronal applications are combinations of TGF-\$\beta\$ with GDNF, both in monomeric form. Also for applications concerning cartilage and/or bone the combination of several monomeric proteins might be useful, like MP52 with a protein of TGF- β (see e.g. WO 92/09697) or MP52 with a cartilage maintenance-inducing protein such as BMP-9 (see e.g. WO 96/39170). When a nucleic acid or an expression vector is used, however, it has to be ensured that when administering to the patient there has to be an - 16 -

environment in which the nucleic acid and the expression vector, respectively, can be expressed and the protein according to the invention can be produced in vivo at the site of action. The same applies accordingly to the host cell according to the invention. When using expression vectors or host cells it is also possible that they encode more than one monomeric protein of the invention to produce a combination of two or more monomeric proteins.

It is advantageous to both the protein and the nucleic acid or the expression vector or the host cell when they are applied in and/or on a biocompatible matrix. The matrix material can be transplanted into the patient, e.g. surgically, wherein the protein either is effective on the surface of the matrix material or the protein or the DNA encoding the protein can be slowly released from the matrix material and then be effective over a long period of time. Additionally it is possible and advantageous to use a biodegradable matrix material in the pharmaceutical composition, wherein this material preferably dissolves during the protein induced tissue formation so that a protein or a nucleic acid contained therein is released and the newly formed tissue replaces the matrix material.

20

15

5

10

Finally, in case of applications relating to bone formation, it is advantageous to use a matrix material which is itself e.g. osteogenically active. By using such a matrix material it becomes possible to achieve a synergistic effect of protein and matrix material and to effect a particularly rapid and effective bone formation.

25

An especially preferred matrix material that can be used according to the invention is a matrix material as described in U.S. 5,231,169 and U.S. 5,776,193 and especially for applications like spinal fusion.

30

When using a combination of a matrix material and protein and/or nucleic acid and/or expression vector, it is preferable to sterilize such a combination

10

15

20

25

30

- 17 -

prior to its use. The matrix and the morphogenetic protein can be separately sterilized and then combined, but it is preferred to terminally sterilize the device consisting of matrix and morphogenetic protein. Terminal sterilization can be achieved by ionizing radation as already described for dimeric proteins (U.S. 5,674,292) but it is also advantageous to use ethylene oxide.

Of course this invention also comprises pharmaceutical compositions containing further substances like e.g. pharmacologically acceptable auxiliary and carrier substances. However, the protein according to the invention, also in case a matrix material is used, does not necessarily have to be used together with this matrix material, but can also be administered systemically, wherein it concentrates preferably in the surrounding of an implanted matrix material.

For some applications the protein according to the invention and the nucleic acid forming this protein, respectively or the expression vector or host cell can preferably be present in an injectable composition. Implants are not necessary or possible for every form of application of the proteins according to the invention. However, it is also possible to provide an implantable vessel or an implantable micropump containing for example semipermeable membranes in which the protein according to the invention or the nucleic acid generating it is contained, from which either one is slowly released over a prolonged period of time. The pharmaceutical composition according to the invention can also contain other vehicles which make it possible that the proteins or the nucleic acids or the expression vectors encoding these proteins be transported to the site of activity and released there, wherein e.g. liposomes or nanospheres can be used. In principle, it is also possible to apply host cells, like e.g. implanted embryonic cells expressing the proteins. Cells transfected with recombinant DNA may be encapusled prior to implantation. Any other practicable but herein not explicitly described form of application of the pharmaceutical composition according the invention and their corresponding manufacture are also comprised by the

10

15

20

25

30

- 18 -

present invention, as long as they contain a protein according to the invention or a nucleic acid or an expression vector coding therefor, or a host cell expressing it.

Although the indications shall not be restricted herein and all indications exhibiting the dimeric form of the protein according to the invention are also comprised, in the following types of application for the compositions according to the invention are listed which are considered to be particularly preferred indications for the proteins of the present invention. On the one hand, there is the prevention or therapy of diseases associated with bone and/or cartilage damage or affecting bone and/or cartilage disease, or generally situations, in which cartilage and/or bone formation is desirable or for spinal fusion, and on the other hand, there is prevention or therapy of damaged or diseased tissue associated with connective tissue including tendon and/or ligament, periodontal or dental tissue including dental implants, neural tissue including CNS tissue and neuropathological situations, tissue of the sensory system, liver, pancreas, cardiac, blood vessel, renal, uterine and thyroid tissue, skin, mucous membranes, endothelium, epithelium, for promotion or induction of nerve growth, tissue regeneration, angiogenesis, wound healing including ulcers, burns, injuries or skin grafts, induction of proliferation of progenitor cells or bone marrow cells, for maintenance of a state of proliferation or differentiation for treatment or preservation of tissue or cells for organ or tissue transplantation, for integrity of gastrointestinal lining, for treatment of disturbances in fertility, contraception or pregnancy.

Diseases concerning sensory organs like the eye are also to be included in the preferred indication of the pharmaceutical composition according to the invention. As neuronal diseases again Parkinson's and Alzheimer's diseases can be mentioned as examples.

10

15

20

25 '

30

- 19 -

The pharmaceutical compositions according to the invention can be used in any desired way, the pharmaceutical compositions are formulated preferably for surgical local application, topical or systemic application. Auxiliary substances for the individual application form can of course be present in the pharmaceutical composition according to the invention. For some applications it can be advantageous to add some further substances to the pharmaceutical composition as for example Vitamin D (WO 92/21365), parathyroid hormone related peptide (WO 97/35607), chordin (WO 98/21335), anti-fibrinolytic agent (EP 535091), anti-metabolites (WO 95/09004), alkyl cellulose (WO 93/00050), mannitol (WO 98/33514), sugar, glycine, glutamic acid hydrochloride (U.S. 5,385,887), antibiotics, antiseptics, amino acids and/or additives which improve the solubility or stablility of the monomeric morphogenetic protein as for example nonionic detergents (e.g. Tween 80), basic amino acids, carrier proteins (e.g. serum albumin), full length propeptides of the TGF-β superfamily or parts thereof.

As can be already gathered from the description of proteins, nucleic acids and pharmaceutical compositions, the proteins according to the invention and respective nucleic acids, which provide for an expression of the proteins at the site of activity, can advantageously be applied in all areas for which also the dimeric forms of the proteins, as described, can be applied. In the framework of the present invention therefore a further subject matter is the use of a pharmaceutical composition according to the present invention for the treatment or prevention of any indications of the dimeric forms of the proteins according to the invention.

Herein it is again possible to conduct surgical operations and to implant the pharmaceutical composition (in particular contained on a matrix material), an administration in liquid or otherwise suitable form via, e.g. injection or oral administration seems to be as suitable as a topical application for e.g. tissue regeneration.

- 20 -

Fig. 1A shows a two dimensional graph of the conformation of recombinantly produced dimeric MP52 with the deleted first alanine. In this figure the 7 cysteine bridges contained in a dimer are shown, wherein there are 3 intramolecular cystine bridges per monomer unit and 1 intermolecular cystine brigde connecting both monomers. Fig. 1B shows the monomeric protein according to the invention wherein the cysteine of the amino acid sequence of MP52 has been replaced by X that denotes any amino acid except cysteine.

Printed:24-08-2000 2:

I IIS PAGE BLANK (USPTO)

- 21 -

SEQUENCE LISTING

```
<110> HyGene AG
5
     <120> Monomeric Protein of the TGF-beta Family
     <130> 20780PEP Monomeric TGF-beta protein
10
     <140>
     <141>
     <160> 4
15
     <170> PatentIn Ver. 2.1
     <210> 1
     <211> 2703
     <212> DNA
20
     <213> Homo sapiens
     <220>
     <221> CDS
25
     <222> (640)..(2142)
     <400> 1
     ccatggcctc gaaagggcag cggtgatttt tttcacataa atatatcgca cttaaatgag 60
30
     tttagacagc atgacatcag agagtaatta aattggtttg ggttggaatt ccgtttccaa 120
     ttcctgagtt caggtttgta aaagattttt ctgagcacct gcaggcctgt gagtgtgtgt 180
35
     gtgtgtgtgt gtgtgtgt gtgtgtgta agtattttca ctggaaagga ttcaaaacta 240
     gggggaaaaa aaaactggag cacacaggca gcattacgcc attcttcctt cttggaaaaa 300
40
     teceteagee ttatacaage etectteaag eeeteagtea gttgtgcagg agaaaggggg 360
```

- 22 -

	cggttggctt tctcctttca agaacgagtt attttcagct gctgactgga gacggtgcac 4	20
5	gtctggatac gagagcattt ccactatggg actggataca aacacacac cggcagactt 4	180
	caagagtete agaetgagga gaaageettt eettetgetg etaetgetge tgeegetget 5	540
10	tttgaaagtc cacteettte atggttttte etgecaaace agaggeacet ttgetgetge	600
	cgctgttctc tttggtgtca ttcagcggct ggccagagg atg aga ctc ccc aaa	654
15	Met Arg Leu Pro Lys 1 5	
	ctc ctc act ttc ttg ctt tgg tac ctg gct tgg ctg gac ctg gaa ttc	702
20	Leu Leu Thr Phe Leu Leu Trp Tyr Leu Ala Trp Leu Asp Leu Glu Phe 10 15 20	
	atc tgc act gtg ttg ggt gcc cct gac ttg ggc cag aga ccc cag ggg	750
25	Ile Cys Thr Val Leu Gly Ala Pro Asp Leu Gly Gln Arg Pro Gln Gly 25 30 35	
	ace agg cca gga ttg gcc aaa gca gag gcc aag gag agg ccc ccc	798
30	Thr Arg Pro Gly Leu Ala Lys Ala Glu Ala Lys Glu Arg Pro Pro Leu 40 45 50	
	gcc cgg aac gtc ttc agg cca ggg ggt cac agc tat ggt ggg ggg gcc	846
35	Ala Arg Asn Val Phe Arg Pro Gly Gly His Ser Tyr Gly Gly Gly Ala 55 60 65	
	acc aat gcc aat gcc agg gca aag gga ggc acc ggg cag aca gga gg	894
40	Thr Asn Ala Asn Ala Arg Ala Lys Gly Gly Thr Gly Gln Thr Gly Gly 70 75 80 85	
	ctg aca cag ccc aag aag gat gaa ccc aaa aag ctg ccc ccc aga ccg	942
	The Cla Dro Lye Lye Asp Glu Pro Lys Lys Leu Pro Pro Arg Pro	

- 23 -

					90					95					100)	
	ggc	ggc	cct	gaa	ccc	aag	cca	gga	cac	cct	ccc	caa	aca	agg	cag	gct	990
5	Gly	Gly	Pro	Glu 105	Pro	Lys	Pro	Gly	His 110	Pro	Pro	Gln	Thr	Arg		Ala	
	aca	gcc	cgg	act	gtg	acc	cca	aaa	gga	cag	ctt	ccc	gga	ggc a	aag	gca	1038
10	Thr	Ala	Arg 120	Thr	Val	Thr	Pro	Lys 125	Gly	Gln	Leu	.Pro	Gly 130		Lys	Ala	
	ccc	cca	aaa	gca	gga	tct	gtc	ccc	agc	tcc	ttc	ctg	ctg	aag a	aag (gcc	1086
15	Pro	Pro 135	Lys	Ala	Gly	Ser	Val 140	Pro	Ser	Ser	Phe	Leu 145		Lys	Lys	Ala	
	agg	gag	ccc	999	ccc	cca	cga	gag	ccc	aag	gag	ccg	ttt	cgc (cca ·	ccc	1134
20	Arg 150	Glu	Pro	Gly	Pro	Pro 155	Arg	Glu	Pro	Lys	Glu 160	Pro	Phe	Arg	Pro	Pro 165	
	ccc	atc	aca	ccc	cac	gag	tac	atg	ctc	tcg	ctg	tac	agg	acg (ctg	tcc	1182
25	Pro	Ile	Thr	Pro	His 170	Glu	Tyr	Met	Leu	Ser 175		Tyr	Arg	Thr	Leu 180	Ser	
	gat	gct	gac	aga	aag	gga	ggc	aac	agc	agc	gtg	aag	ttg	gag (gct :	ggc	1230
30	Asp	Ala	Asp	Arg 185	Lys	Gly	Gly	Asn	Ser 190	Ser	Val	Lys	Leu	Glu 195	Ala	Gly	
	ctg	gcc	aac	acc	atc	acc	agc	ttt	att	gac	aaa	3 33	caa	gat q	gac	cga	1278
35	Leu	Ala	Asn 200	Thr	Ile	Thr	Ser	Phe 205	Ile	Asp	Lys	Gly	Gln 210		Asp	Arg	
	ggt	ccc	gtg	gtc	agg	aag	cag	agg	tac	gtg	ttt	gac	att	agt 9	gcc ·	ctg	1326
40	Gly	Pro 215	Val	Val	Arg	Lys	Gln 220	Arg	Tyr	Val	Phe	Asp 225		Ser	Ala	Leu	
	gag	aag	gat	a aa	ctg	ctg	999	gcc	gag	ctg	cgg	atc	ttg	cgg (aag	aag	1374
45	Glu	Tave	Δsn	Glv	Len	T.en	Glv	Ala	Glu	Leu	Arg	Ile	Leu	Ara	Lvs	Lvs	

- 24 -

	230					235					240					245	
	ccc	tcg	gac	acg	gcc	aag	cca ·	gcg	gcc	ccc (gga ç	ggc g	ggg (egg s	jct g	cc	1422
5	Pro	Ser	Asp	Thr	Ala 250	Lys	Pro	Ala	Ala	Pro 255	Gly	Gly	Gly	Arg	Ala 260	Ala	
	cag	ctg	aag	ctg	tcc	agc	tgc	ccc	agc	ggc	cgg (cag (eeg g	gee t	cc t	tg	1470
10	Gln	Leu	Lys	Leu 265	Ser	Ser	Cys	Pro	Ser 270	Gly	Arg	Gln	Pro	Ala 275	Ser	Leu	
	ctg	gat	gtg	cgc	tcc	gtg	cca	ggc	ctg	gac	gga 1	tct s	ggc 1	tgg g	gag g	tg	1518
15	Leu	Asp	Val 280	Arg	Ser	Val	Pro	Gly 285	Leu	Asp	Gly	Ser	Gly 290	Trp	Glu	Val	
	ttc	gac	atc	tgg	aag	ctc	ttc	cga	aac	ttt	aag	aac 1	tcg :	gac (cag c	tg	1566
20	Phe	Asp 295		Trp	Lys	Leu	Phe 300	Arg	Asn	Phe	Lys	Asn 305	Ser	Ala	Gln	Leu	
	tgc	ctg	gag	ctg	gag	gcc	tgg	gaa	cgg	ggc	agg :	gcc (gtg (gac (ctc c	gt	1614
25	Cys 310	Leu	Glu	Leu	Glu	Ala 315		Glu	Arg	Gly	Arg 320		Val	Asp	Leu	Arg 325.	
	ggc	ctg	ggc	ttc	gac	cgc	gcc	gcc	cgg	cag	gtc	cac	gag	aag g	gcc c	tg	1662
30	Gly	Leu	Gly	Phe	Asp 330		Ala	Ala	Arg	Gln 335		His	Glu	Lys	Ala 340	Leu	
•	ttc	ctg	gtg	ttt	ggc	cgc	acc	aag	aaa	cgg	gac	ctg	ttc	ttt	aat g	ag	1710
35	Phe	Leu	Val	Phe		Arg	Thr	Lys	150 350		Asp	Leu	. Phe	Phe 355	Asn	Glu	
	att	aag	gcc	cgc	tct	ggc	cag	gac	gat	aag	acc	gtg	tat	gag	tac c	tg	1758
40	Ile	Lys	360		Ser	Gly	Gln	Asp 365		Lys	s Thr	· Val	. Туг 370		Tyr	Leu	
	ttc	ago	cag	cgg	cga	aaa	cgg	cgg	gcc	cca	ctg	gcc	act	cgc	cag s	gc	1806
4 5	Dhe	Ser	r Gln	Arc	r Arc	LVS	. Arc	Arc	r Ala	a Pro	Lev	ı Ala	Thi	Arc	Gln	Gly	

- 25 -

		375					380	ı				385	i					
	aag	cga	ccc	agc	aag	aac	ctt	aag	gct	cgc	tgc	agt	cgg	aag	gca	ctg	1854	1
5	Lys 390	Arg	Pro	Ser	Lys	Asn 395	Leu	Lys	Ala	Arg	Cys 400		Arg	Lys	s Ala	405		
	cat	gtc	aac	ttc	aag	gac	atg	ggc	tgg	gac	gac	tgg	atc	atc	gca	ccc	1902	2
10	His	Val	Asn	Phe	Lys 410	Asp	Met	Gly	Trp	415		Trp	Ile	Ile	420	a Pro		
	ctt	gag	tac	gag	gct	ttc	cac	tgc	gag	aaa	ctg	tgc	gag	ttc	cca	ttg	1950)
15	Leu	Glu	Tyr	Glu 425	Ala	Phe	His	Cys	Glu 430	_	Leu	Cys	Gl u	Phe 435) Leu		
	cgc	tcc	cac	ctg	gag	ccc	acg	aat	cat	gca	gtc	atc	cag	acc	ctg	atg	1998	3
20	Arg	Ser	His 440	Leu	Glu	Pro	Thr	Asn 445		Ala	Val	Ile	Gln 450		Lev	ı Met		
•	aac	tcc	atg	gac	ccc	gag	tcc	aca	cca	ccc	acc	nnn	tgt	gtg	ccc	acg	2046	
25	Asn	Ser 455	Met	Asp	Pro	Glu	Ser 460		Pro	Pro	Thr	Xaa 465		Val	. Pro	Thr		
	cgg	ctg	agt	ccc	atc	agc	atc	ctc	ttc	att	gac	tct	gcc	aac	aac	gtg	2094	ŀ
30	Arg 470	Leu	Ser	Pro	Ile	Ser 475	Ile	Leu	Phe	lle	Asp 480		Ala	Asn	Asr	val 485		
	gtg	tat	aag	cag	tat	gag	gac	atg	gtc	gtg	gag	tcg	tgt	ggc	tgc	agg	2142	:
35	Val	Tyr	Lys	Gln	Tyr 490	Glu	Asp	Met	Val	Val 495		Ser	Cys	Gly	500	Arg		
	tago	cagca	act g	gcco	etate	gt ct	tect	egggt	ggo	cacat	ccc	aaga	gccc	ct t	cctg	cacte	2202	:
40	ctgg	gaato	cac a	ıgagg	ggto	ca gg	Jaago	etgte	g gca	ıggag	rcat	ctac	acag	ct t	gggt	gaaag	2262	:
45	333	attco	caa t	aago	ettge	et cg	rctcl	tctga	a gtg	gtgac	ttg	ggct	aaag	gc c	ccct	tttat	2322	:

- 26 -

	ccacaa	agtt	c cc	ctgg	ctga	ggat	tgct	tgc (ccgtc	tgct	g at	gtga	ccag	tgg	cagg	cac :	2382
5	aggtco	cagg	g ag	acag	actc	tga	atgg	gac 1	cgagt	ccca	ıg ga	aaca	gtgc	ttt	ccga	tga :	2442
	gactc	agco	c ac	catt	tctc	ctc	acct	ggg (cctto	ctcag	ge et	ctgg	gacto	tcc	taag	cac	2502
10	ctctc	agga	ıg ag	ccac	aggt	gcc	actg	cct	cctca	aaato	ca ca	tttg	rtged	tgg	rtgac	ttc	2562
	ctgtc	ccts	ag ga	ıcagt	tgag	, aag	ctga	.ctg	ggca	agagi	tg gg	gagag	gaaga	a gga	ıgagg	gct	2622
15	tggat	agag	gt t <u>s</u>	jagga	agtgt	gag	gctg	rtta	gact	gtta	ga t	ttaa	atgta	a tat	tgat	gag	2682
20	ataaa	ıaage	ca a	aact	gtgc	c t											2703
25	<210 × <211 × <212 × <213 × <213 × <213 × <	> 50 > PR	T	apie	ns												
30	<400: Met 1	Arg			5					10					15		
	Leu i	Asp	Leu	Glu 20	Phe	Ile	Cys	Thr	Val 25	Leu	Gly	Ala	Pro	Asp 30	Leu	GIA	
35	Gln .	Arg	Pro 35	Gln	Gly	Thr	Arg	Pro 40	Gly	Leu	Ala	Lys	Ala 45	Glu	Ala	Lys	
	Glu	Arg 50	Pro	Pro	Leu	Ala	Arg 55	Asn	Val	Phe	Arg	Pro 60	Gly	Gly	His	Ser	
40	Tyr 65	Gly	Gly	Gly	Ala	Thr 70		Ala	Asn	Ala	Arg 75		Lys	Gly	Gly	Thr 80	
45	Gly	Gln	Thr	Gly	Gly 85		Thr	Gln	Pro	Lys 90		Asp	Glu	Pro	Lys 95	Lys	

6

- 27 -

									2,							
	Leu	Pro	Pro	Arg 100	Pro	Gly	Gly	Pro	Glu 105	Pro	Lys	Pro	Gly	His 110	Pro	Pro
5	Gln	Thr	Arg 115	Gln	Ala	Thr	Ala	Arg 120	Thr	Val	Thr	Pro	Lys 125	Gly	Gln	Leu
	Pro	Gly 130	Gly	Lys	Ala	Pro	Pro 135	Lys	Ala	Gly	Ser	Val 140	Pro	Ser	Ser	Phe
10	Leu 145	Leu	Lys	Lys	Ala	Arg 150	Glu	Pro	Gly	Pro	Pro 155	Arg	Glu	Pro	Lys	Glu 160
15	Pro	Phe	Arg	Pro	Pro 165	Pro	Ile	Thr	Pro	His 170	Glu	Tyr	Met	Leu	Ser 175	Leu
	Tyr	Arg	Thr	Leu 180	Ser	Asp	Ala	Asp	Arg 185	Lys	Gly	Gly	Asn	Ser 190	Ser	Val
20	Lys	Leu	Glu 195	Ala	Gly	Leu	Ala	Asn 200	Thr	Ile	Thr	Ser	Phe 205	Ile	Asp	Lys
	Gly	Gln 210	Asp	Asp	Arg	Gly	Pro 215	Val	Val	Arg	Lys	Gln 220	Arg	Tyr	Val	Phe
25	Asp 225	Ile	Ser	Ala	Leu	Glu 230	Lys	Asp	Gly	Leu	Leu 235	Gly	Ala	Glu	Leu	Arg 240
30	Ile	Leu	Arg	Lys	Lys 245	Pro	Ser	Asp	Thr	Ala 250	Lys	Pro	Ala	Ala	Pro 255	Gly
	Gly	Gly	Arg	Ala 260	Ala	Gln	Leu	Lys	Leu 265	Ser	Ser	Cys	Pro	Ser 270	Gly	Arg
35	Gln	Pro	Ala 275	Ser	Leu	Leu	Asp	Val 280	Arg	Ser	Val	Pro	Gly 285	Leu	Asp	Gly
	Ser	Gly 290	Trp	Glu	Val	Phe	Asp 295	Ile	Trp	Lys	Leu	Phe 300	Arg	Asn	Phe	Lys
40	Asn 305	Ser	Ala	Gln	Leu	Cys 310	Leu	Glu	Leu	Glu	Ala 315	Trp	Glu	Arg	Gly	Arg 320
	Ala	Val	Asp	Leu	Arg 325	Gly	Leu	Gly	Phe	Asp 330	Arg	Ala	Ala	Arg	Gln 335	Val

- 28 -

His Glu Lys Ala Leu Phe Leu Val Phe Gly Arg Thr Lys Lys Arg Asp 340 345 350

Leu Phe Phe Asn Glu Ile Lys Ala Arg Ser Gly Gln Asp Asp Lys Thr 355 360 365

Val Tyr Glu Tyr Leu Phe Ser Gln Arg Arg Lys Arg Arg Ala Pro Leu 370 375 380

Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala Arg Cys 385 390 395 400

Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp Asp Asp 405 410 415

Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly Leu 420 425 430

Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His Ala Val 20 435 440 445

Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro Thr 450 455 460

25 Xaa Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile Asp 465 470 475 480

Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val Glu 485 490 495

Ser Cys Gly Cys Arg

<210> 3

<211> 2272

<212> DNA

<213> Homo sapiens

40

30

35

<220>

<221> CDS

<222> (128)..(1183)

45 <400> 3

- 29 -

	caag	ggag	cca	tgcc	agct	gg a	caca	cacti	t ct	tcca	gggc	ctc	tggc	agc	cago	racagag	60
5	ttga	agac	cac a	agcto	gttga	ag ac	cacto	gagco	c ctg	gagto	etgt	atto	gata	aag a	aaggg	geette	120
	ccca	agca	atg	acc	tcc	tca	ttg	ctt	ctg	gcc	ttt	ctc	ctc	ctg	gct	cca	169
10			Met 1	Thr	Ser	Ser	Leu 5	Leu	Leu	Ala	Phe	Leu 10	Leu	Leu	Ala	Pro	
	acc	aca	gtg	gcc	act	ccc	.aga ·	gct	ggc	ggt	cag	tgt	cca	gca	tgt	a aa	217
15	Thr 15	Thr	Val	Ala	Thr	Pro 20	Arg	Ala	Gly	Gly	Gln 25	Cys	Pro	Ala	Cys	Gly 30	
	3 33	ccc	acc	ttg	gaa	ctg	gag	agc	cag	cgg	gag	ctg	ctt	ctt	gat	ctg	265
20	Gly	Pro	Thr	Leu	Glu 35	Leu	Glu	Ser	Gln	Arg 40	Glu	Leu	Leu	Leu	Asp 45	Leu	
	gcc	aag	aga	agc	atc	ttg	gac	aag	ctg	cac	ctc	acc	cag	cgc	cca	aca	313
25	Ala	Lys	Arg	Ser 50	Ile	Leu	Asp	Lys	Leu 55	His	Leu	Thr	Gln	Arg 60	Pro	Thr	
	ctg	aac	cgc	cct	gtg	tcc	aga	gct	gct	ttg	agg	act	gca	ctg	cag	cac	361
30	Leu	Asn	Arg 65	Pro	Val	Ser	Arg	Ala 70	Ala	Leu	Arg	Thr	Ala 75	Leu	Gln	His	
	ctc	cac	3 33	gtc	cca	cag	999	gca	ctt	cta	gag	gac	aac	agg	gaa	cag	409
35	Leu	His 80	Gly	Val	Pro	Gln	Gly 85	Ala	Leu	Leu	Glu	Asp 90	Asn	Arg	Glu	Gln	
	gaa	tgt	gaa	atc	atc	agc	ttt	gct	gag	aca	ggc	ctc	tcc	acc	atc	aac	457
40	Glu 95	Cys	Glu	Ile	Ile	Ser 100	Phe	Ala	Glu	Thr	Gly 105	Leu	Ser	Thr	Ile	Asn 110	
	cag	act	cgt	ctt	gat	ttt	cac	ttc	tcc	tct	gat	aga	act	gct	ggt	gac	505
4E	Gln	Thr	Arg	Leu	Asp	Phe	His	Phe	Ser	Ser	Asp	Arg	Thr	Ala	Gly	Asp	

- 30 -

	agg	gag	gtc	cag	cag	gcc	agt	ctc	atg	ttc	ttt	gtg	cag	ctc	cct	tcc	553
	Arg	Glu	Val	Gln 130	Gln	Ala	Ser	Leu	Met 135	Phe	Phe	Val	Gln	Leu 140	Pro	Ser	
5	aat	acc	act	tgg	acc	ttg	aaa	gtg	aga	gtc	ctt	gtg	ctg	ggt	cca	cat	601
	Asn	Thr	Thr 145	Trp	Thr	Leu	Lys	Val 150	Arg	Val	Leu	Val	Leu 155	Gly	Pro	His	
10	aat	acc	aac	ctc	acc	ttg	gct	act	caq	tac	ctg	ctg	gag	gtg	gat	gcc	649
						_										Ala	
	ASII	160	ASII	neu	1111	neu	165		GIII	171	Deu	170	GLU	val	rap	ALG	
15	agt	ggc	tgg	cat	caa	ctc	ccc	cta	333	cct	gaa	gct	caa	gct	gcc	tgc	697
	Ser 175	Gly	Trp	His	Gln	Leu 180	Pro	Leu	Gly	Pro	Glu 185	Ala	Gln	Ala	Ala	Cys 190	
20	agc	cag	aaa	cac	ctg	acc	ctg	gag	ctg	gta	ctt	gaa	ggc	cag	gta	gcc	745
	Ser	Gln	Gly	His	Leu 195	Thr	Leu	Glu	Leu	Val 200	Leu	Glu	Gly	Gln	Val 205	Ala	
25	cag	agc	tca	gtc	atc	ctg	ggt	gga	gct	gcc	cat	agg	cct	ttt	gtg	gca	793
	Gln	Ser	Ser	Val 210	Ile	Leu	Gly	Gly	Ala 215	Ala	His	Arg	Pro	Phe 220	Val	Ala	
30	gcc	cgg	gtg	aga	gtt	ggg	ggc	aaa	cac	cag	att	cac	cga	cga	ggc	atc	841
	Ala	Arg	Val 225	Arg	Val	Gly	Gly	Lys 230	His	Gln	Ile	His	Arg 235	Arg	Gly	Ile	
35	gac	tgc	caa	gga	3 33	tcc	agg	atg	tgc	tgt	cga	caa	gag	ttt	ttt	gtg	889
40	Asp	Cys 240	Gln	Gly	Gly	Ser	Arg 245	Met	Cys	Cys	Arg	Gln 250	Glu	Phe	Phe	Val	
-10	gac	ttc	cgt	gag	att	ggc	tgg	cac	gac	tgg	atc	atc	cag	cct	gag	ggc	937
AE	Asp 255	Phe	Arg	Glu	Ile	Gly 260	Trp	His	Asp	Trp	Ile 265		Gln	Pro	Glu	Gly 270	

- 31 -

	tac	gcc	atg	aac	ttc	tgc	ata	a aa	cag	tgc	cca	cta	cac	ata	gca	ggc	985
_	Tyr	Ala	Met	Asn	Phe 275	Cys	Ile	Gly	Gln	Cys 280	Pro	Leu	His	Ile	Ala 285	Gly	
5	atg	cct	ggt	att	gct	gcc	tcc	ttt	cac	act :	gca :	gtg (ctc a	aat (ctt (ctc	1033
10	Met	Pro	Gly	Ile 290	Ala	Ala	Ser	Phe	His 295	Thr	Ala	Val	Leu	Asn 300	Leu	Leu	
	aag	gcc	aac	aca	gct	gca	ggc	acc	act 9	gga g	ggg (ggc 1	ca r	nnn t	gt (gta	1081
15	Lys	Ala	Asn 305	Thr	Ala	Ala	Gly	Thr 310	Thr	Gly	Gly	Gly	Ser 315	Xaa	Cys	Val	
15	ccc	acg	gcc	cgg	cgc	ccc	ctg	tct	ctg (ctc 1	tat 1	tat q	gac a	agg g	gac a	gc	1129
20	Pro	Thr 320	Ala	Arg	Arg	Pro	Leu 325	Ser	Leu	Leu	Tyr	Tyr 330	Asp	Arg	Asp	Ser	
20	aac	att	gtc	aag	act	gac	ata	cct	gac a	atg g	gta g	gta g	gag g	jcc t	gt g	iaa	1177
25	Asn 335	Ile	Val	Lys	Thr	Asp 340	Ile	Pro	Asp	Met	Val 345	Val	Glu	Ala	Cys	Gly 350	
25	tgc	agt	tagt	ctat	gt g	tggt	atgg	g ca	gccc	aagg	ttg	catg	gga a	aaaca	acgc	cc	1233
	Cys	Ser															
30	ctac	agaa	ıgt g	rcact	tect	t ga	gagg	aggg	aatg	gacci	tca t	tete	etgto	c ag	_l aat <u>ç</u>	ıtgga	1293
	ctcc	ctct	tc c	tgag	catc	t ta	tgga	aatt	acco	ccaco	ctt t	gact	tgaa	ıg aa	acct	tcat:	1353
35	ctaa	agca	ag t	cact	gtgc	c at	cttc	ctga	ccad	ctaco	cct c	ettto	cctag	ig gc	atag	rtcca	1413
40	tccc	gcta	ıgt c	cato	ccgc	t ag	cccc	actc	cagg	ggact	tca ç	gacco	atct	c ca	acca	ıtgag	1473
•	caat	gcca	ıtc t	ggtt	ccca	g gc	aaag	acac	cctt	agct	tca c	ettt	aata	ıg ac	ccca	taac	1533

	ccactatgcc	ttcctgtcct	ttctactcaa	tggtccccac	tccaagatga	gttgacacaa	1593
5	cccttccc	caatttttgt	ggatctccag	agaggccctt	ctttggattc	accaaagttt	1653
	agatcactgc	tgcccaaaat	agaggettae	ctaccccct	ctttgttgtg	agcccctgtc	1713
10	cttcttagtt	gtccaggtga	actactaaag	ctctctttgc	ataccttcat	ccattttttg	1773
	tccttctctg	cctttctcta	tgcccttaag	gggtgacttg	cctgagctct	atcacctgag	1833
15	ctcccctgcc	ctctggcttc	ctgctgaggt	cagggcattt	cttatccctg	ttecetetet	1893
20	gtctaggtgt	catggttctg	tgtaactgtg	gctattctgt	gtccctacac	tacctggcta	1953
	ccccttcca	tggccccagc	tctgcctaca	ttctgatttt	tttttttt	tttttttga	2013
25	aaagttaaaa	attccttaat	tttttattcc	tggtaccact	accacaattt	acagggcaat	2073
	atacctgatg	taatgaaaag	aaaaagaaaa	agacaaagct	acaacagata	aaagacctca	2133
30	ggaatgtaca	tctaattgac	actacattgc	attaatcaat	agctgcactt	tttgcaaact	2193
35	gtggctatga	cagtcctgaa	caagaagggt	ttcctgttta	agctgcagta	acttttctga	2253
	ctatggatca	tegtteett					2272
40							
	<210> 4 <211> 352						

•

45

<212> PRT

<213> Homo sapiens

- 33 -

	/4U	U - 4														
	Met 1	Thr	Ser	Ser	Leu 5	Leu	Leu	Ala	Phe	Leu 10	Leu	Leu	Ala	Pro	Thr 15	Th
5	Val	Ala	Thr	Pro 20	Arg	Ala	Gly	Gly	Gln 25	Cys	Pro	Ala	Cys	Gly 30	Gly	Pro
40	Thr	Leu	Glu 35	Leu	Glu	Ser	Gln	Arg 40	Glu	Leu	Leu	Leu	Asp 45	Leu	Ala	Ly
10	Arg	Ser 50	Ile	Leu	Asp	Lys	Leu 55	His	Leu	Thr	Gln	Arg 60	Pro	Thr	Leu	Ası
15	Arg 65	Pro	Val	Ser	Arg	Ala 70	Ala	Leu	Arg	Thr	Ala 75	Leu	Gln	His	Leu	His
	Gly	V al	Pro	Gln	Gly 85	Ala	Leu	Leu	Glu	Asp 90	Asn	Arg	Glu	Gln	Glu 95	Суя
20	Glu	Ile	Ile	Ser 100	Phe	Ala	Glu	Thr	Gly 105	Leu	Ser	Thr	Ile	Asn 110	Gln	Thi
25	Arg	Leu	Asp 115	Phe	His	Phe	Ser	Ser 120	Asp	Arg	Thr	Ala	Gly 125	Asp	Arg	Glu
	Val	Gln 130	Gln	Ala	Ser	Leu	Met 135	Phe	Phe	Val	Gln	Leu 140	Pro	Ser	Asn	Thi
30	Thr 145	Trp	Thr	Leu	Lys	Val 150	Arg	Val	Leu	Val	Leu 155	Gly	Pro	His	Asn	Th:
•	Asn	Leu	Thr	Leu	Ala 165	Thr	Gln	Tyr	Leu	Leu 170	Glu	Val	Asp	Ala	Ser 175	Gly
35	Trp	His	Gln	Leu 180	Pro	Leu	Gly	Pro	Glu 185	Ala	Gln	Ala	Ala	Cys 190	Ser	Glr
40	Gly	His	Leu 195	Thr	Leu	Glu	Leu	Val 200	Leu	Glu	Gly	Gln	Val 205	Ala	Gln	Ser
	Ser	Val 210	Ile	Leu	Gly	Gly	Ala 215	Ala	His	Arg	Pro	Phe 220	Val	Ala	Ala	Arg
A E		Arg	Val	Gly	Gly	_	His	Gln	Ile	His	Arg 235	Arg	Gly	Ile	Asp	Cys
45	225					230					233					~ * (

- 34 -

Gln Gly Gly Ser Arg Met Cys Cys Arg Gln Glu Phe Phe Val Asp Phe 245 250 255

Arg Glu Ile Gly Trp His Asp Trp Ile Ile Gln Pro Glu Gly Tyr Ala 5 260 265 270

Met Asn Phe Cys Ile Gly Gln Cys Pro Leu His Ile Ala Gly Met Pro 275 280 285

10 Gly Ile Ala Ala Ser Phe His Thr Ala Val Leu Asn Leu Leu Lys Ala 290 295 300

Asn Thr Ala Ala Gly Thr Thr Gly Gly Gly Ser Xaa Cys Val Pro Thr 305 310 315 320

Ala Arg Arg Pro Leu Ser Leu Leu Tyr Tyr Asp Arg Asp Ser Asn Ile
325 330 335

Val Lys Thr Asp Ile Pro Asp Met Val Val Glu Ala Cys Gly Cys Ser 340 345 350

5

10

15

20

25

- 35 -

EPO-Munich 52

0 6. Aug. 1999

Claims

- Protein selected from the members of the TGF-ß superfamily, characterized in that the protein is necessarily monomeric due to substitution or deletion of a cysteine which is responsible for dimer formation.
- Protein according to claim 1, characterized in that the protein is a mature protein or a biologically active part or variant thereof.
- Protein according to any one of the preceding claims, characterized in that the protein contains at least the 7 cysteine region characteristic for the TGF-ß protein superfamily.
- 4. Protein according to claim 3, characterized in that it contains a consensus sequence according to Formula I: C(Y)₂₅₋₂₉CYYYC(Y)₂₅₋₃₅XC(Y)₂₇₋₃₄CYC or Formula II: C(Y)₂₈CYYYC(Y)₃₀₋₃₂XC(Y)₃₁CYC, wherein C denotes cysteine, Y denotes any amino acid and X denotes any amino acid except cysteine.
- Protein according to any one of claims 1 to 4,
 characterized in that the protein is a morphogenetic protein.
- 6. Protein according to any one of the preceding claims, characterized in that the proteins belongs to the TGF-β, BMP, GDF, activin or GDNF family.
- 7. Protein according to claim 6, characterized in that the protein is MP52 (GDF5) or a biologically active part or variant thereof.

- 36 -

- 8. Protein according to any one of the preceeding claims, characterized in that it comprises the amino acid sequence according to SEQ.ID.NO.2 or a part thereof.
- 9. Protein according to claim 6, characterized in that the protein is MP121 or a biologically active part or variant thereof.
- 10. Protein according to claim 9,

 characterized in that it comprises the amino acid sequence according to SEQ.ID.NO.4 or a part thereof.
 - 11. Protein according to any one of claims 1 to 10, characterized in that the cysteine residue is substituted by an amino acid selected from the group of alanine, serine, threonine, leucine, isoleucine, glycine and valine.
- 12. Protein according to any one of claims 1 to 11,
 characterized in that it contains additional amino acids that
 facilitate or mediate the transfer and localization of the protein in a
 certain tissue.
 - 13. Nucleic acid, characterized in that it encodes a protein according to any one of claims 1 to 12.
 - 14. Nucleic acid according to claim 13, characterized in that it is a DNA.
- 30 15. Nucleic acid according to claim 13 or 14, characterized in that it contains a sequence as shown in SEQ.ID.NO.1 or a fragment thereof.

15

- 37 -

- 16. Nucleic acid according to claim 13 or 14, characterized in that it contains a sequence as shown in SEQ.ID.NO.3 or a fragment thereof.
- 5 17. Nucleic acid according to any one of claims 13 to 16, characterized in that it further contains suitable expression control sequences facilitating and/or driving expression of the encoded protein.
- 10 18. Expression vector, characterized in that it contains a nucleic acid according to any one of claims 13 to 17 in a suitable vector system.
- 19. Expression vector according to claim 18,characterized in that the vector system is suitable for prokaryotic expression.
- 20. Host cell,
 characterized in that it contains a nucleic acid according to any
 one of claims 13 to 17 or an expression vector according to claims
 18 or 19 and upon expression of said nucleic acid or vector is able
 to produce a monomeric protein according to any one of claims 1
 to 12.
- 25 21. Host cell according to claim 20, characterized in that it is a prokaryotic host cell.
 - 22. Host cell according to claim 20, characterized in that it is an embryonal cell.
 - 23. Pharmaceutical composition,

5

10

15

20

25

- 38 -

characterized in that it contains at least one protein according to any one of claims 1 to 12 or at least one nucleic acid according to any one of claims 13 to 17, at least one expression vector according to any one of claims 18 or 19 or at least one host cell according to claim 20 or 22.

- 24. Pharmaceutical composition according to claim 23, characterized in that the protein and/or nucleic acid are contained in and/or on a biocompatible matrix material.
- 25. Pharmaceutical composition according to claim 24, characterized in that the matrix material is biodegradable.
- 26. Pharmaceutical composition according to claims 24 or 25, characterized in that the matrix material is itself osteogenically active.
- 27. Pharmaceutical compostion according to any one of claims 23 to 26, for the prevention or therapy of diseases for which also the dimeric form of the protein would be indicated.
 - 28. Pharmaceutical composition according to claim 27, for prevention or therapy of diseases associated with bone and/or cartilage damage or affecting bone and/or cartilage disease or situations in which cartilage and/or bone growth is desirable or for spinal fusion.
- 29. Pharmaceutical composition according to claim 27,
 for prevention or therapy of damaged or diseased tissue associated
 with connective tissue including tendon and/or ligament,
 periodontal or dental tissue including dental implants, neural tissue

- 39 -

including CNS tissue and neuropahtological situations, tissue of the sensory system, liver, pancreas, cardiac, blood vessel, renal, uterine and thyroid tissue, skin, mucous membranes, endothelium, epithelium, for promotion or induction of nerve growth, tissue regeneration, angiogenesis, wound healing including ulcers, burns, injuries or skin grafts, induction of proliferation of progenitor cells or bone marrow cells, for maintenance of a state of proliferation or differentiation, for treatment or preservation of tissue or cells for organ or tissue transplantation, for integrity of gastrointestinal lining, for treatment of disturbances in fertility, contraception or pregnancy.

- 30. Pharmaceutical composition according to any one of claims 23 to 29 for surgical local application, topical or systemic application.
- 31. Pharmaceutical compostion according to any one of claims 23 to 30 characterized in that it further contains pharmacologically acceptable auxiliary substances.
- 32. Pharmaceutical composition according to any one of claims 30 or31,characterized in that the composition is injectable.
- 25 33. Pharmaceutical composition according to anyone of claims 30 to 32, characterized in that it is contained in a vehicle that allows to direct and release the composition to a determined site of action.
- 30 34. Pharmaceutical composition according to claim 33, characterized in that the vehicle is selected from liposomes, nanospheres, larger implantable containers and micropumps.

. . 13

06-08-1999

5

10

15

- 40 -

- 35. Use of a pharmaceutical composition according to any one of claims 23 to 34 for the prevention or treatment of any indications of the dimeric form of the protein.
- ₅ bb/20780PEP 04.08.1999

- 41 -

EPO-Munich 52

06, Aug. 1999

Summary

The present invention is concerned with proteins selected from the members of the TGF-ß superfamily, which are monomeric due to substitution or deletion of a cysteine which is responsible for dimer formation.

The invention is also concerned with nucleic acids, encoding such monomeric proteins, vectors or host cells containing the nucleic acids as well as with pharmaceutical compositions comprising the proteins or nucleic acids encoding the proteins. The pharmaceutical compositions can be applied advantageously for all indications for which the respective dimeric proteins are useful.

15

5

10

cl 19.07.1999



Ex. 1

Fig. 1 A

EPO-Munich 52

0 6. Aug. 1999

Name: MP52, dimeric form

Formula

 $C_{1184}H_{1844}N_{330}O_{350}S_{22}$

Molecular weight

26994 Dalton

Amino acid composition

238 amino acids

Disulfide bond

7 bonds

Primary structure

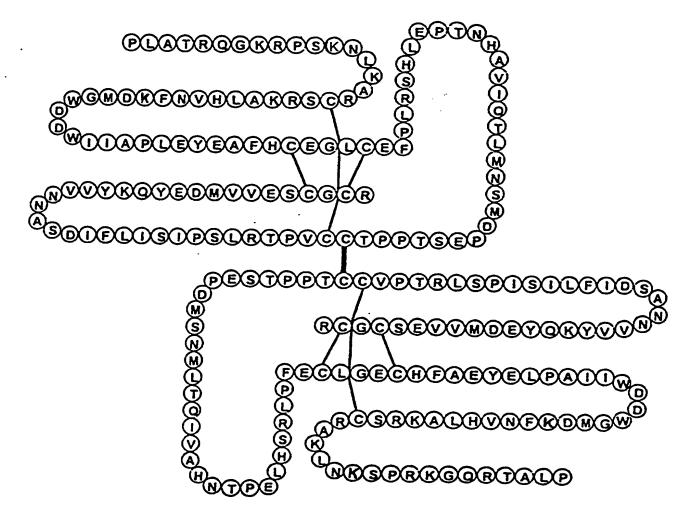


Fig. 1 B

